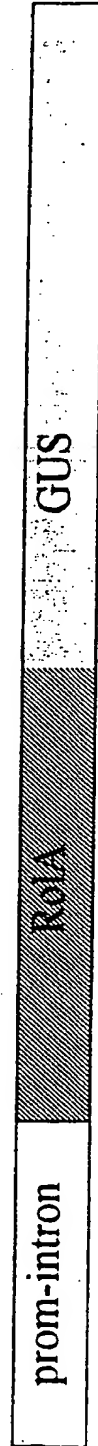


PROMOTER REGION REPORTER GENE CONSTRUCTS



86AGUS



85-17 GUS



Δ GUS

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FIG. 1

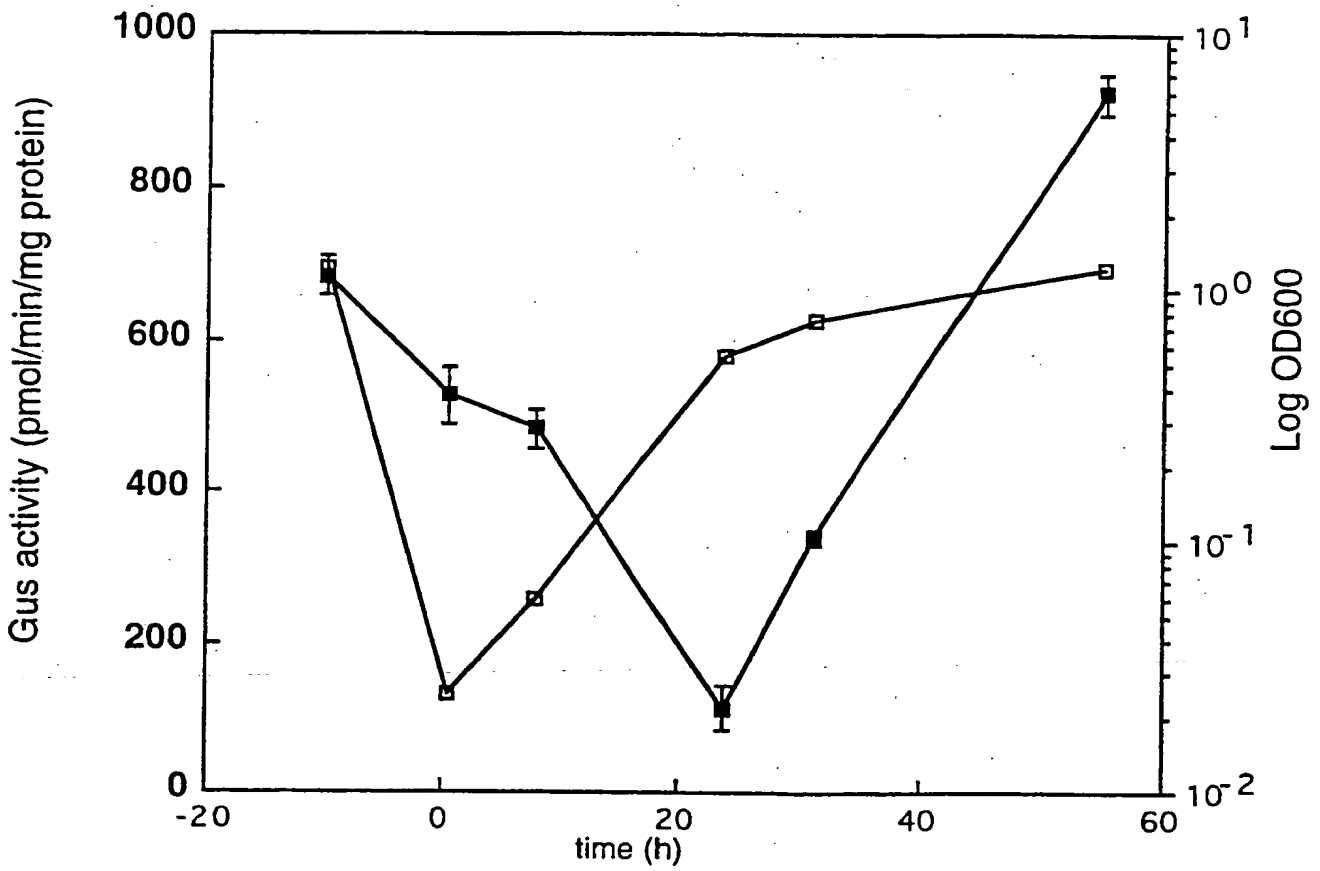


FIG. 2

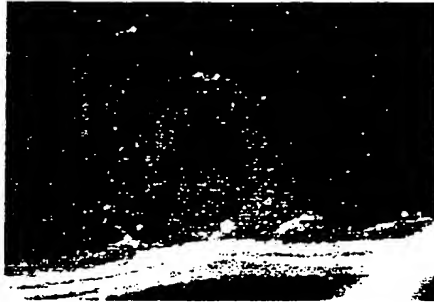
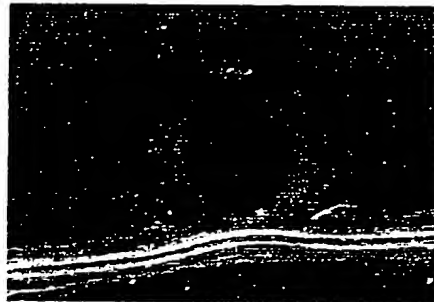
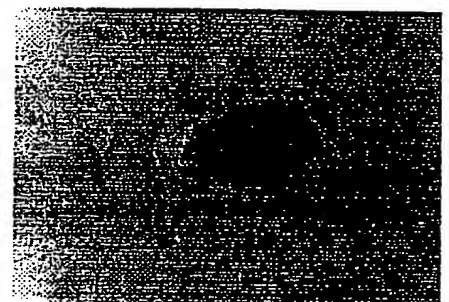
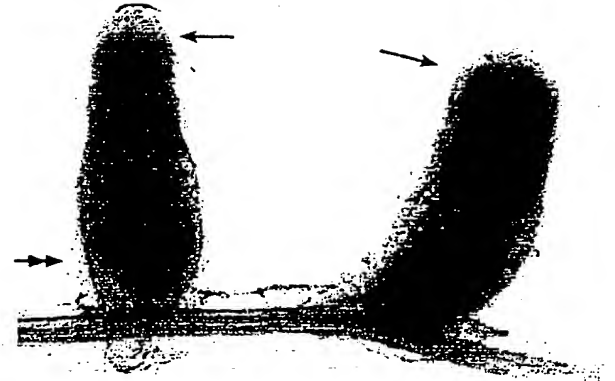
**A****B****C****D****E****F****G****H**

FIG. 3

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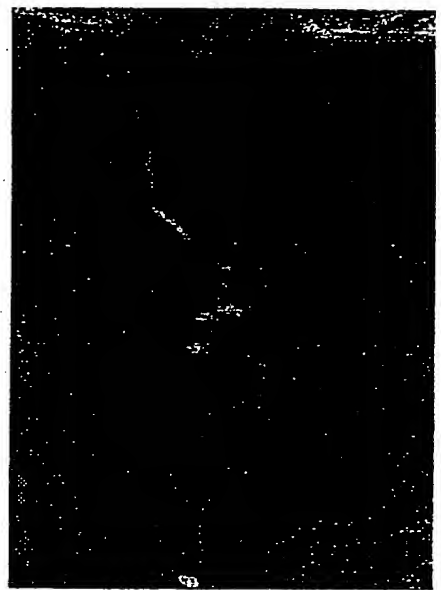
FIG. 4



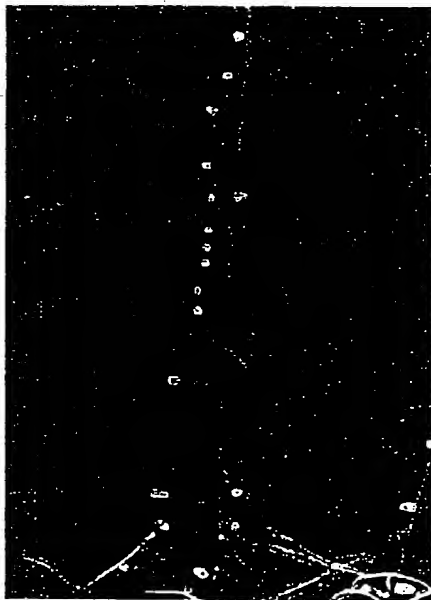
A



B



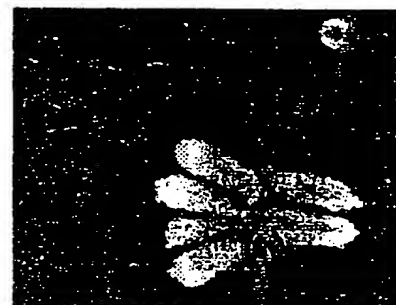
C



D



E1



E



F



G

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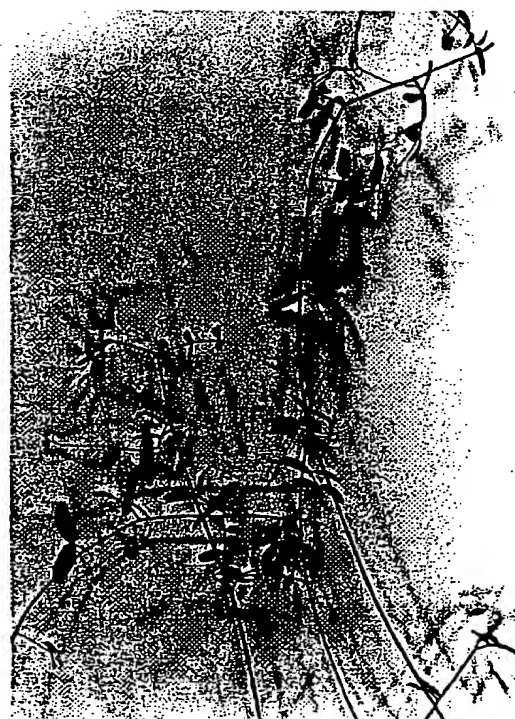
**A****B****C****D**

FIG. 5

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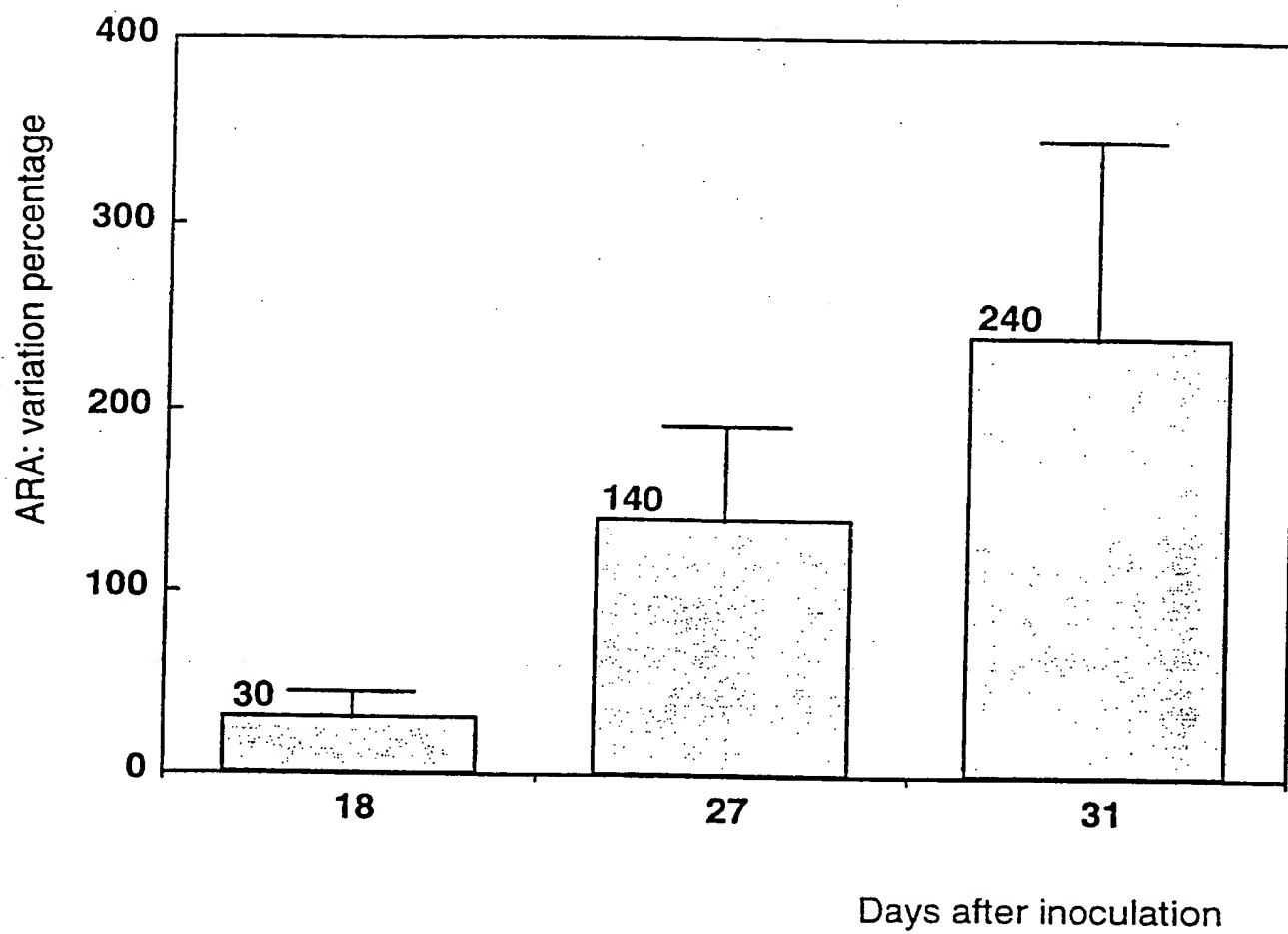


FIG. 6

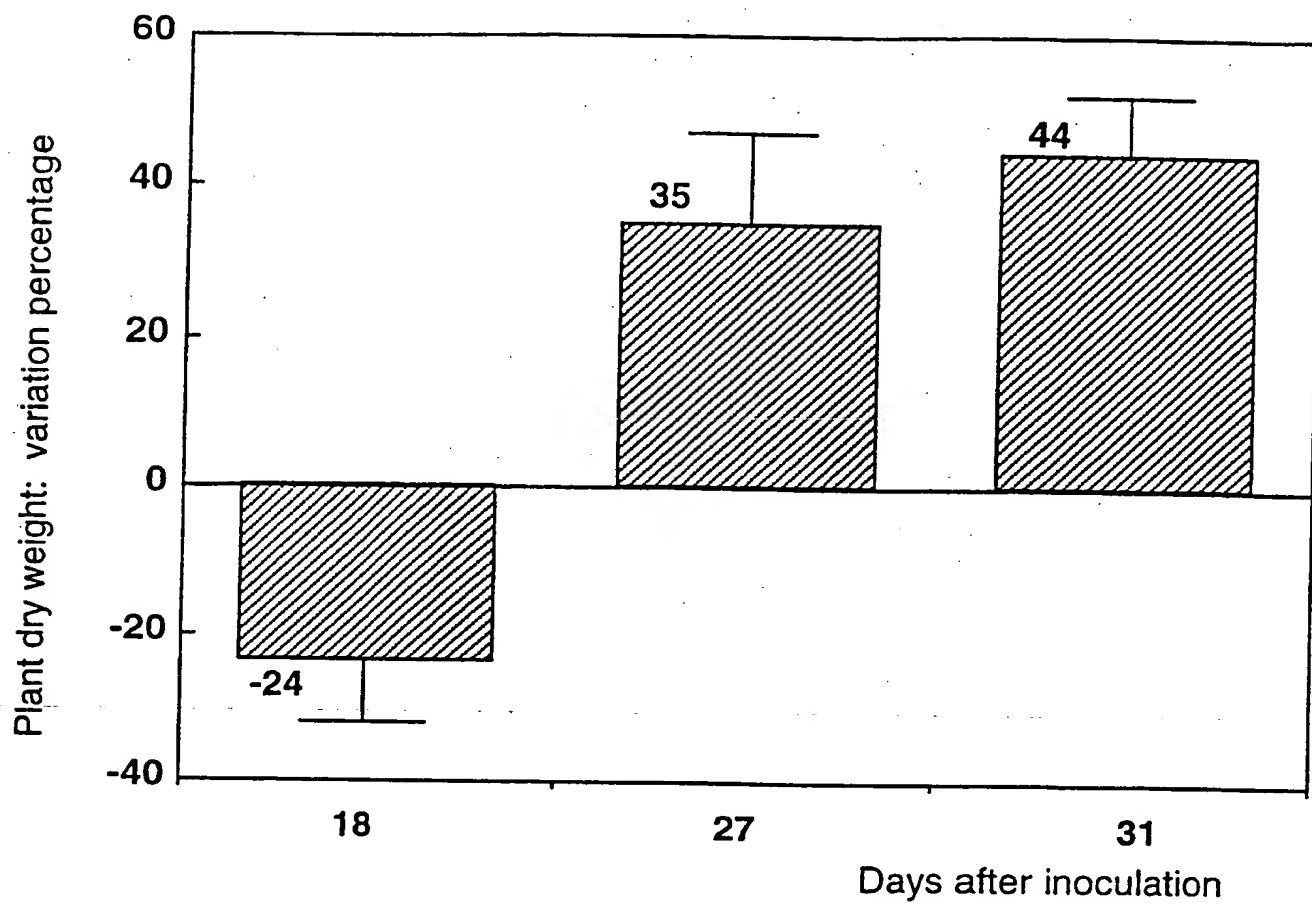


FIG. 7

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+1

└→

s.s. GTGAGTGTGGTTGTAGGTTCAATTATTACTATTTTGAAGCTGTGTATT-TCCCTTTTCTAAATATGCACCTATTTCATGTTTCAGAAATG// roIA A4 coding region
s.s.

TTGAAT____17bp____TATAAT
-35 -10

σ⁷⁰ CONSENSUS
SEQUENCE

CTATACT

-10

σ³⁸ CONSENSUS
SEQUENCE

CGGCAAGT

GEAR BOX
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FIG. 8

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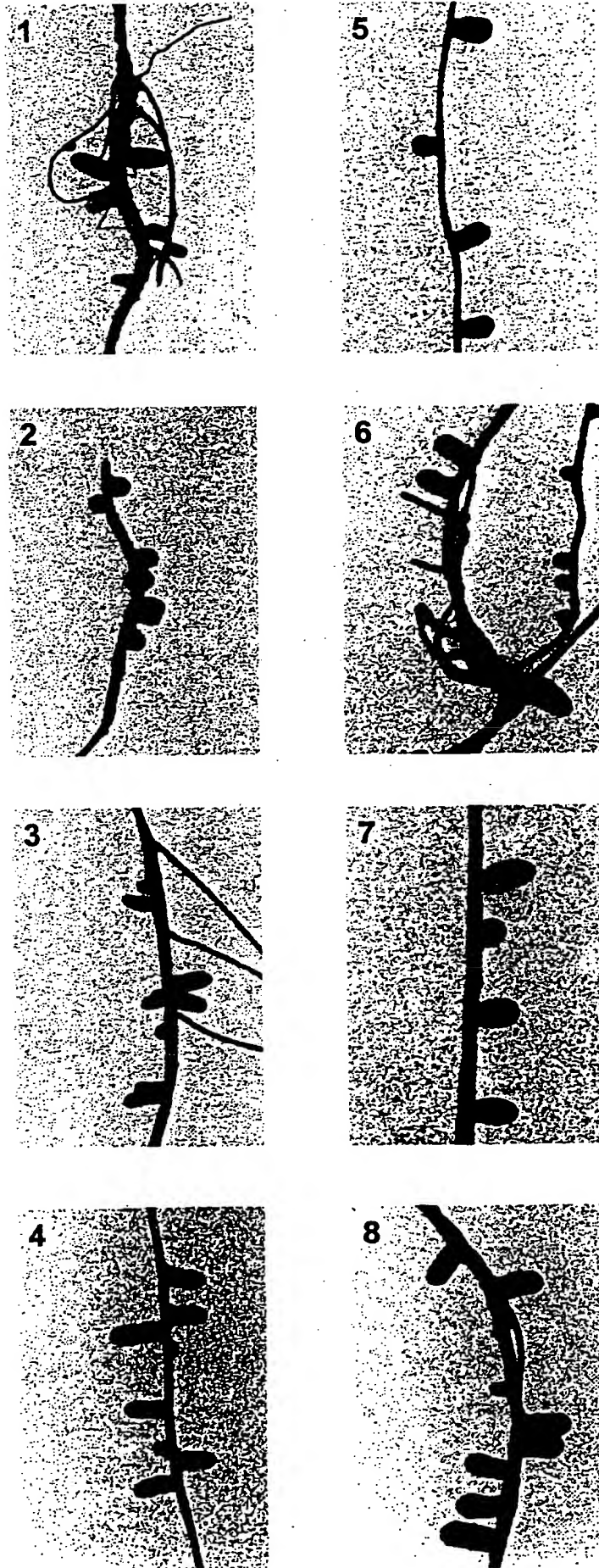


FIG. 9

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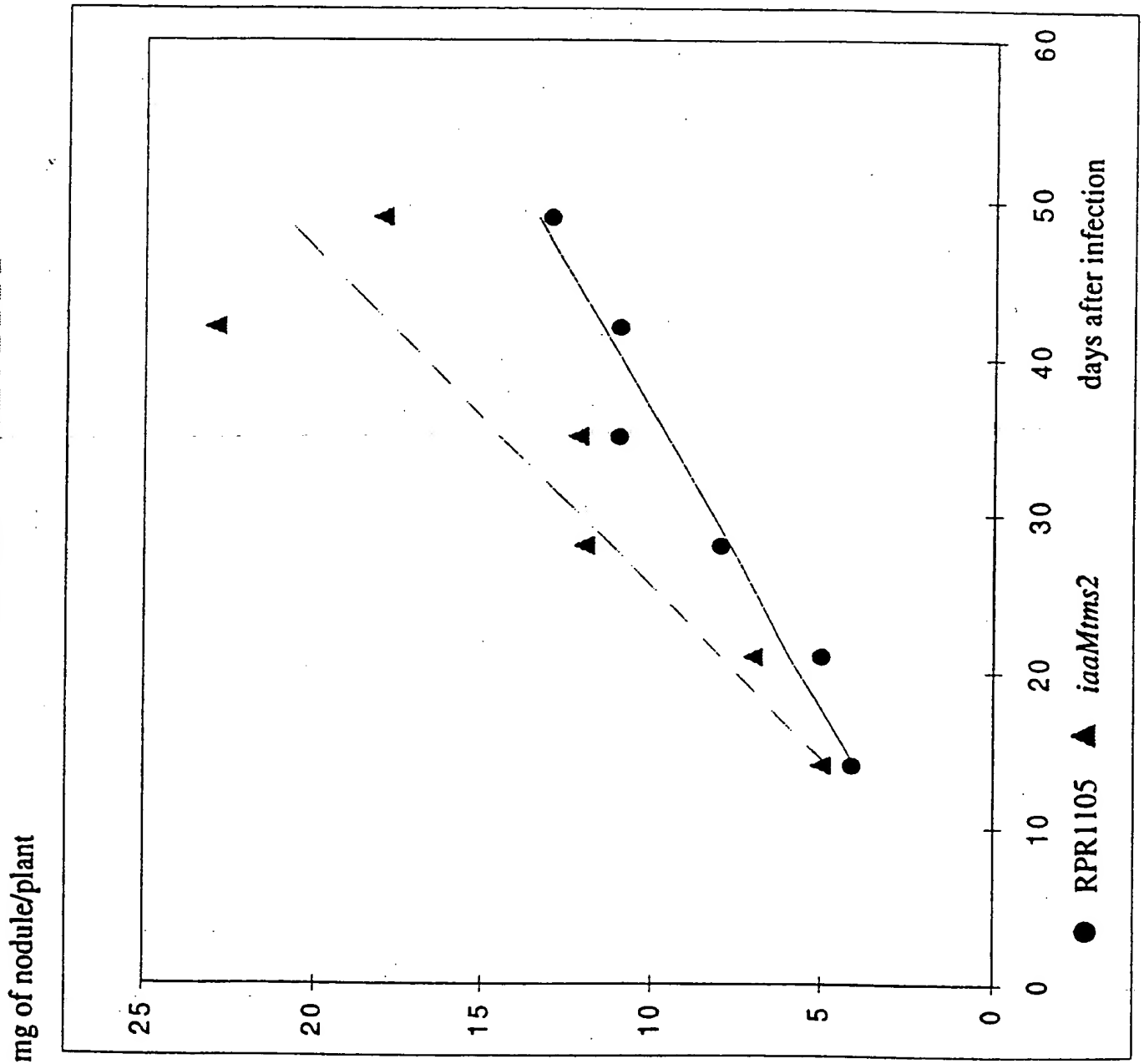
Days A. I.	Mnod(mg)/pp	
	RPR 1105	<i>iaaMtms2</i>
14	4 ± 1	5,0 ± 0,4
21	5 ± 1	7 ± 2
28	8 ± 2	12 ± 3
35	11 ± 1	12,2 ± 0,3
42	11 ± 4	23 ± 6
49	13 ± 3	18 ± 4

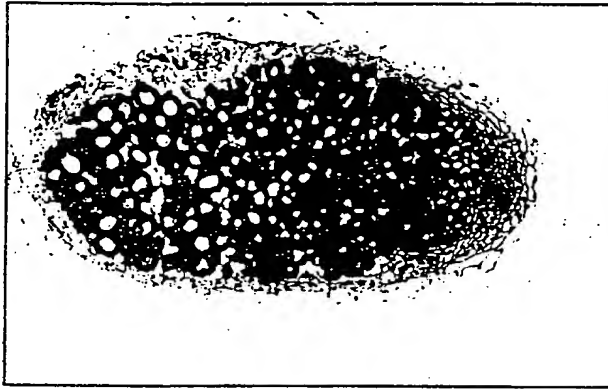
pendence	RPR 1105	0,26700	±	3,E-05
intercept	RPR 1105	0,26	±	0,03

pendence	<i>iaaMtms2</i>	0,4620	±	1,E-04
intercept	<i>iaaMtms2</i>	-1,69	±	0,1

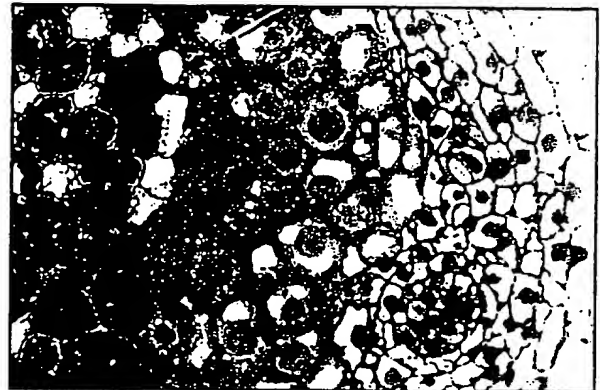
FIG. 10A

FIG. 10B





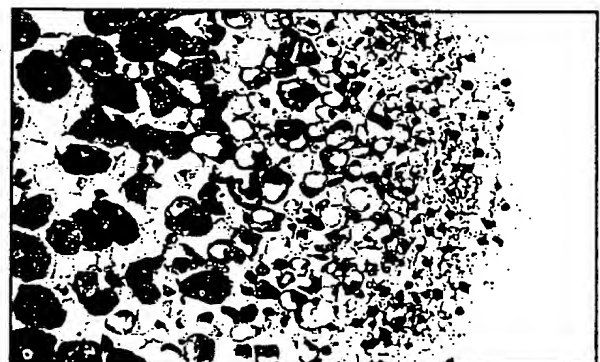
1



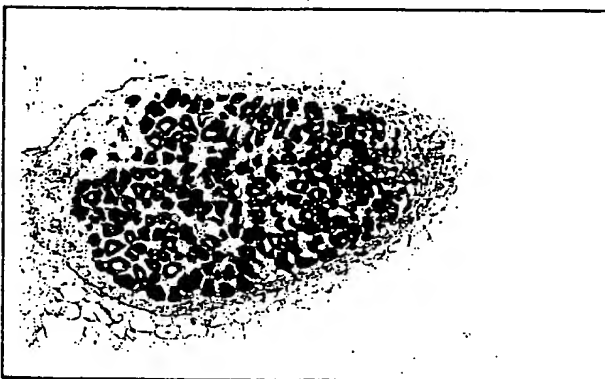
2



3



4



5

FIG. 11

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